

SEQ ID NO:16

Rat Smooth Muscle Myosin Heavy Chain Gene Sequence (-4,216 to +11,795)

Nucleotide 1 corresponds to -4,216 bp relative to the SM-MHC transcription start site

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	CAGGGTCCCT	CACTGAACCT	GTTGGAGCTA	TGCTGGTAGC	CAGCAAGGCC	CAGTGGCCTT	180
	CCTGTCCTCA	TCTCACACAG	CACAATATGT	GTGGCCATGC	TCCACTTTTT	TACATGGAAA	240
	TTGGGGTCTT	CCAACTGGGG	TTCTCATTTG	TGCAGTGACA	CTCTTCCCCA	CTGAGGCCATC	300
	TCCTCAGGCC	AGCTGTATATA	TTTTAAATA	ATTAATATT	TAGCACATGC	CTTTAGAAGC	360
	CAATAGCTAT	TTAAAGCTGT	TTGCTTAAAAA	AAAAAAAAAA	AAAAAAAGACT	TCATTATCCC	420
	AACACTTATG	AGGGAGAGAC	AATAATTCCA	AAACCAGAAC	CAGCCAGGGT	ACACAGTGAG	480
	ACTTTATTTA	AAAAAAA	AAAAGAAAG	AAAGAAAAAA	AAAAGAAAAA	GAAAAAAA	540
	GGCTCCAAAG	AGAAATTTC	CCTTCATCAT	CTAATCACAA	GAAAACAATT	TATTTATTT	600
	GACATCACTC	AGTCCAAAGG	AGCTTTTGT	AAAGTGACTT	CTCTTCTTAA	AATAAGTGAC	660
	CCTTCCCAAC	CACCAAAAC	AAAACAGAAA	CCTCTGCCCT	GTTCTAGAGT	CCTTTGAAG	720
	ACTTCAGATA	CCTGAAGAGT	GGACAGATAT	TTACCGAGTG	ACTTAAATGA	ACATACTGTC	780
	CCTGGTACT	GCTCAAGCAT	GCCAGGAGAG	CATGGATGGT	TTATGCAAGG	CTGGCACTGT	840
	CATTAACAAC	TCAGTAAGGC	GGAGAAGACA	GAGAGCCTCT	CCTAAGACAA	TGGCACATAA	900
	GGACATGGGT	AAACCCAGAG	GTTCCCGGCT	AGTACTTAGC	AGAGCTGAGA	TCAGACTTGG	960
	GCCTCTGTGC	TCGCTTGCT	AGTGGGCAAC	ACTCAAGACT	GGGGTAAACA	ATAAGTTGAT	1020
	CTGGGATATG	GCTCAGTAAT	CACACTGAGA	ATTCAACACT	GGGAAGGCAG	AGGAGGATCC	1080
	CTGGGATTGC	TGCCTGGCTC	TCTAGCAGCC	TAGCAGAATC	AACAAACTCC	AGGTTAGTGTG	1140
	AGAGATGCTC	ACAAAATAAA	ATGGAGGAGC	AACTGAACAC	ACTCAGTGTGTT	GACCCACACA	1200
	CACACTAAAG	AACACGTGTA	CCACACAGAC	ACAGACACAG	GATAACCTAC	CCATGTTGTG	1260
	TATGGACTCA	GCCAGCCCCAG	GTTGGAAACT	CAGTTCCCT	GTTAACTCTT	TTCAAACCTG	1320
	GGTCCTCAGC	GATGTGCTGG	GGAACCTACT	TCACGGCATT	ATTCTGGGCA	TTAGATGTAA	1380
	AGGAAGCAGT	AAAGTTTCCC	TTTCTTGAC	TGAGGTGATG	CGAGAATGAG	GGCCTGAATT	1440
	CCATCTCTAG	GACTCACATA	AAGACACCCA	GACTGCACTG	GCCAGTAAGC	CTCACCTATG	1500
	CCTCCAAGCC	TGGCTGTGAG	AGACTGTCTC	AAAAACAAAG	TAAAACAAAC	AAAATCAATG	1560
	TCAGATGTGC	ACACATCGAA	TCCCAGCATG	TGTACGGCAT	GCTTGCAGTC	AGCCTTGTTT	1620
	ACAGAGAGTT	CTAGGCCAAC	CAGCTATACA	CAGTGAGACC	CTGTGGTAGA	CGGCTCCTAA	1680
	GAAC TGACAT	TTGTGACTGA	CAGATGTGCA	CATCTACCAC	ATGCACATCA	CAGTTCCAT	1740
	TTTACAAAAA	GGTTAACACT	TACTAATTGA	TTAGGGAGTG	GGGCACCCCA	CTGCTACATG	1800
	TGAAAGCCAG	AGAATGATGT	GTTCCAGTCG	GTCAGTTGTG	TCCTTCCACC	ATGTAGGTCC	1860
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	TCAGCTGTGC	TCCTTGGAGT	TTGACTTCAC	TGAAGCCTGC	TACAGGAGTG	CCCTTCCCTC	2040
	CTAGCACTAG	GATGGCCAGC	TCTGGGCTGG	TTTCAGACTA	GGGTAGGTGC	AGGTGGGCC	2100
	TGGGCTTCCC	TCCCTCATTC	CTCCTGGGCT	CAATGCCAAG	CCGGTTTCCA	TTCCCTTTAC	2160
	GTGCACTGCG	AAGAGGCTT	GGGGAAAGCGG	CCTCATCCAT	CATGCAGAGA	GCTCCCTCCC	2220
	CACCTCTACA	GAGAGCCAGC	CAAGCTGCTG	TCCTTGGCTC	TGCTCTGTCC	ACCCCTGTGAG	2280
	GAGGCTGGGA	TGAGGTTGGG	GATGGGGAGG	ATCAGGATTG	AGATGTTTC	AAGTCTGAGA	2340
	AGCAGGTGAG	CTTGGTCTTA	GAAGAATATG	GAAGGGGTCT	ACTGGGGTTG	AGATATAGAT	2400
	CACTGTATCA	AAAGTCACAG	GGGGGCTGTG	TGGCTTTTTC	ATATCCAAA	GTCAGCTTGG	2460
	TGCTGTTTC	CTAGGCTTCC	TGAGTCCGAC	AAAGGTGCAG	TGTGTTAATC	TCACACCACT	2520
	TCAAGGACTG	TTACAAAAAA	AAAATAGGA	GGAGCTCGAT	TCGCCCCTTT	TTACAGGCAG	2580
	GGTAACTAAG	AGCCAGTACT	TGCCCATGGT	CCTGCTGTTA	TAAAGAGGCT	CAGTAGACTC	2640
	CCATTCAAAC	AACTGTGCTC	AGAGGCCCTC	TGTCGTCCTG	TGGCCAATTG	CCCTATTGCT	2700
	CTCTGGAGTG	AATATTGGGA	TATTAACAG	TACTGACCTT	GCTGAGGACC	CTCAGGGTAC	2760
	TCAGCTCTTC	TGGCCTGCAA	AATGGGGCTG	GGACAGGTTG	GCCAGGATCA	TCCTCTGGTT	2820
	GGGAGAACCA	GCTGCACGTG	GGTCTGGAGC	TCTTATTAGT	ACTGGGGTCC	CCATAACGCT	2880
	CCATGGGCTC	AGCAGGGAGGC	TGCAACGGGAC	CATATTAGT	CAGGGGGAGC	CAGAGCCCCG	2940
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	TCCCAGCCAG	GCCCAGGAGG	GAGGACCAGC	TCAGGACCTC	GAGGGTCCGT	GGCGGGGGAG	3060
	CGAGGCGTCC	CCGGCCTGGC	ATGAGGCCAA	CTCTGCTCG	ACTTCCTTTT	ATGGCCTGAG	3120

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	GAGAGACACA	CAGAGAGAGA	CAGAGAGACA	GAGAGACACA	CAGAGAGAGA	CAGAGACAGA	3300
	CACACACAGA	GAGAGACAGA	CAGACAAAGA	GAGAGACAGA	GACAGAGAGA	CACACACAGA	3360
	GAGACAGACA	GACAAAAAGA	GAAGAGAGAC	AGAGACTTTA	GGGACGTAAT	CATCACAGGG	3420
5	AAATCAAAGC	TAAGAGTGTG	ATGAAAAGAG	TGTCAGGTCA	GACAAAAGAG	ACAGGGGCCA	3480
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	TGACAGCAGG	TCCCCCACAT	TCTCTTAGAG	TCTTAGCATG	CATCCTCCAA	GTGCCATAAC	3600
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10	CCGGTTGTAG	GGTGAAGGA	GCTGCAGAGA	ACAAGTTGGA	AAAACAAGTT	TCCCAGCAGT	3720
	CACAGAGGAT	ATGCAGTGAC	TGTGCCACT	TGTTTTTTT	TTTTTAAGTC	CCCTCCCCC	3780
	CCCCCGCCCC	CCCCCGGCT	TGCTAAGCAC	AACC GGCTTC	GAATCTTAGG	AAGTGGCAGG	3840
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	CAGAGGCTCT	GTTCCTCACCC	TGCAGGGGGC	GCTGGGAAGG	GCAGAGGACC	CTCCCACCC	4140
	GCCCCGCACT	CACCTCCCCC	TCCCCACCC	CGGGTAGCGC	TGACTCTATA	AAGCCAGATG	4200
20	→ transcription start site +1						
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	CTCCATCCGG	TGTTCTCCCTG	CTAGTCCACC	CCAGTAGCAG	ATCTGTAAGT	AGAAGTTGAT	4320
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25	GGTTGCCTAT	GAACAGAACCC	ACCTGGGAAA	GTGGGGTAGG	TAATTAAAGG	TTCTGCCAC	4560
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	ACACTGAGTG	AGCAGACGAT	GGATTTGGG	GCTCCTCAGT	GGGAAGGTTA	CTCTCAGGTC	5040
	AGGGAGAGGA	GCTAGCAGAG	AAATTATGC	TATTCCAGTT	CAGAATTGGA	GAAGTCTTGC	5100
35	CATGTCAGA	AAGCACCCCT	CAAAGTTATG	TCTGTCAAGAG	AACAGAAAAA	TTTTTTTGTG	5160
	AAGCCAGGAC	AAGGCTGCTT	TGGTTCTACT	ACTAAGAACT	GAAAAACTGC	TGACTTGCTG	5220
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	GAGCATATTG	TGTATGACAC	AGCTGATATC	AAGAAACCCA	AACGGTGGCC	TTTCCCCTAA	6900
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	GTTGTCAGG	GCAACAAAGC	CTCAGGGAGC	AGCCAGATGA	CCTCACTCCG	TTTTGGCCA	12360
	GAGACACAAA	CTTTGCACTT	GATCTTGT	GTGCTTTAA	GCCCGTTT	AGATGAGGTT	12420
40	CCTGGAAAAG	CTAATCTCA	CGTCTTTCA	TTTTTCTGTT	GAACCTTTCG	TGATGTTTC	12480
	TAACCTAATT	GCAATTAAA	AAGAGGCAGC	TTGCTGTCCA	GGAGGAATGA	CACAAACACT	12540
	AGGCCTCTGA	GTGACTAAAG	ACCATTGAA	ATGGGTCGTC	ATCTATTACA	GAAAATGTAA	12600
	AATATACTTT	ACACTTCTA	ACTATGTGC	TAAAGTATGT	TTTATTTGT	TTTCTCTAA	12660
	AAAAAGAATT	ATTTATTTA	CGTATTGAG	TACACTGTAG	CTGACTTCAG	ATCCACCAAGA	12720
45	AGAGGGCCTT	AGATTCCATT	ACAGATGGTT	GTGAGCTACC	GTGTGATGGG	AATTGAACTC	12780
	AGGACCTCTG	GA AAA AGCAGT	CAGTGTCTT	AACCACTGAG	CCATCTTCC	GGCCTTTATT	12840
	TTCCTTTTTT	TA A A A A A A A A	ATAAATGAA	AATTAACCTT	TATTCATGG	GTGTATATAT	12900
	GTATGGCTC	AAACATGATA	TATGTGCATG	GGCTCACACA	TGCAGTGGTG	CATGTATAAA	12960
	AGTCAGAGAC	AACTTGCA	AGATGGTTG	CTCTTTCTAT	CATATGGGCC	CTGAGGATTA	13020
50	AACTCAAGTC	ATCAGTTTTT	GTGCCAACCC	CCTTTACTCC	CCGAGCCTTC	TCTCAACAGC	13080
	TCCTCACTTT	ACCTTTTTAT	TTAAAAAACAA	AACAAACAA	CAAACACCAA	CCCAGCCTCC	13140
	CACACAACAA	CGAAAAGATC	TCATGTAGCC	CCAGGGTGGC	TTTGAACCTCC	CCATATAGCT	13200
	TAGGATGACT	TTGAATTCTC	AATGTTCTG	CCTCTACCTC	CTAGTTACTA	TGCCTGGCTT	13260
	CTTACCATAG	AATTTAAGAA	ATTATCTAAG	GTAAAGTGGT	GT T AT GT GT	TATAAGCCAG	13320
55	GCACTCAGGA	AGAAGCTAAG	GCATGATGAT	TGTGAGTTG	AAGCCAAACCC	AGGTTACAGA	13380
	GGATCTCATC	AAGAAATCAA	CATTCAATT	TCAATTATTT	CTTAAATT	TTGAGGTTGG	13440
	GCTGGAGGGG	TTGGTTAAGA	GCACTGGTTG	GTCTTCCAGA	GGACATGAGT	TTGATTCCCT	13500
	GTACCCACAA	TGGTGGCTCA	CAACCATCTG	TAATTTAAT	TCTAGGGATC	TAACGCCCTC	13560
	TTCAAGCCTT	CTCAGGCAGG	TGCATAAGTA	CACAGTCATA	CATGCACAGA	AAACACATAA	13620
	ACATAAAATA	AATAAATTAA	AATTGAGAA	GT T T T T T T G	GGTGGAAAGGT	ACTTTTAAGT	13680

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AACATTCTAT	GTTATGGAAC	AAGTGCATTC	AATTTTACTA	AGTTTTTAAT	TTTAGCTTT	13740
TGTTTGTGG	TTTCTGTTT	GGAAACAAGGT	CTTGTGTATC	CCAAGCATCC	TCAAAGTTGT	13800
TGTGTAGCGA	AGGATGACCT	TGAATTTTTT	TATACTACTG	CCTCTTGAG	GGCAAGCATT	13860
TTAATATAGG	CAAAATAAAC	TTTAAACTTT	GTTTGCTGTG	CAGGTATATA	TGGTGTGCAA	13920
GTGTATCTGT	GTGTGTGTG	GTGTGTGTG	GTGTGTGTG	GTGTGTGAGA	GAGAGAGAGA	13980
GAGAGAGAGA	GAGAGAGAGA	GAGAGAGAGA	GATTAGAGAA	TAACCTGTGG	AAGTTCTCTC	14040
CTTCTACCCCT	GTGGGTCCCC	GGGTAAACTC	GGGTTATAAG	GCTTGCACC	CTTTTCCC	14100
CTGAGAACTT	CTTGTGGCC	TCACTCCCTA	TTTTATTGTTA	TTGGTGGCAG	TACTATTGCT	14160
TTTGAATCCC	ATCTGAAGCT	TGTTTTGTT	GTTTGGTTTT	TAAGGCAGTC	TTAACTGTGA	14220
CCTAAGCTGG	TTTAAAACCT	ACAGGAATT	TCCACCTCCA	CCTCCCAAGT	GTTGGGTTA	14280
CAGATGTGAG	CCCCAAGCCT	GAGTGTCTCT	GAAAGCTGCT	TTTTTTTATT	TCAAAACTAT	14340
CTTTCTCTG	TGTGTAGGTC	TGATTAGTTG	TGGGGTTAGG	TGGTGTCA	ATGATCCATC	14400
ACTCTCCAGC	TATTATTCTT	AAAATGAAGG	GTCTGGGGC	TGGGGATT	GCTCAGTGGT	14460
AGAGCGCTTA	CCTAGGAAGC	GCAAGGCCCT	GGGTCGGTC	CCCAGCTCCG	AAAAAAAGAA	14520
CCCCCC	AAAAAATGAA	GGGTCGGTG	GCTGAGGAAA	AAGCTCAGTT	GCAAAAAAAC	14580
ATGAAAACCT	GATTCAATCT	GTAAAGCCCA	CATAAAAGCC	AGGCATGGCG	GCATGCACCT	14640
ATAACCCCAG	CACTGGGAA	ACAGAACAGG	AGAATACCAA	GAACCTGCTG	GTCAGTCAGT	14700
CTAGTTAAT	TGGTGAGCTC	CAAGCTCAGT	GAGACCTGT	CTCAAAATA	AATGGAGATG	14760
ATCTGTCATC	AAGACCTGGC	CTCCATACAT	ATATGCACAC	ATGTTACTCC	CTCACATGAA	14820
ACATATTAT	AAACAAACAT	ATGCACACAC	TTGTGCATAC	ATGAACAGAT	ATCTATATTG	14880
GCATACACAT	AAAAACACAC	ACACACATAT	ATATATACAA	AAGTGTGTAC	AAACATAGGC	14940
ATAGTATACA	ACCATGCATA	AATGCACAGT	CACACATATG	AATGCATTCA	TATTACACACA	15000
TGGACACATG	AACACATACA	TATATGCTAT	ATCTTATATT	ACACTCCATT	ACTATCCCCC	15060
AGTCCAGGTT	TCAAATATT	ACAAACAGAA	AAGCGGGCTA	CTACCTGTAC	TTTTTCCCAA	15120
TTGCCTTGA	ACAGCGATCT	CTCGACACCT	GATCCCCGCA	GTGCTCCCTG	CGGCAGAGCT	15180
TCATCCGGAA	ACAACCCCCA	TGCACTCTAT	TGATTTTAAT	ACTGGGGATT	ACCTGGAGCC	15240
TTGTAAAGCT	AAACACATTG	TCTACTGCTA	AATACTTCAT	TCTTGCCCC	TTTCCCATGG	15300
GGCGTTTCA	ATCCAGTTAT	TTTTAGTGTG	TTCTTAGATT	TAAGCATCCA	CTAGTACAGA	15360
TTCAAGGATA	TTTTTATTAT	CCCCAAATA	ACAGTATTG	TTAGGTGTAA	CCTTGTAGTT	15420
TTTCCCCAGC	GGCTAATT	AATTGCTTTC	ATGAATAGCC	TATTCTGGAA	AAGTAATT	15480
TTTTTTTTT	TTTTTTTTG	GGTTCTTTT	TTCGGAGCTG	GGGACCGAAC	CCAGGGCCTT	15540
GCGCTCCTA	GGTAAGCGCT	CTACCACTGA	GCTAAATCCC	CAGCCCCAAT	TCTGGACATT	15600
TCTTATAAAAT	GTCACTATGC	TGTATGTGTT	CTTTCAGCAT	TGCAACACTT	TGGTTCTTT	15660
TTATGGCTCA	ATACTGGCT	ACTTATGGAT	CTACCACACT	ATCTATCCAT	TCATCTAAC	15720
ATAGTCATGG	GTGGTATTTC	TACTTGGGG	CTATTATAAG	CTTGCTAGGA	GTATTATGA	15780
CCACATCTT	AGATGCACTG	ATGCATT	TTATCCTAAG	AACAGATCCT	GGATCATATG	15840
GTGGTCTGT	GTTCAAACAT	CAGAGGCACC	ACCATTATT	TTATAATAGG	CATTAAAGAT	15900
TTGGGTATCT	TCTAACTGGG	TGGTGGTGGT	ACATGCCTGT	AGTCCCAGCT	CCTGGGAGGC	15960
AGAGGCAAGT	AGATCCGAAT	TCTCGCCCTA	TAGTGAGTCG	TATTAGTCGA	C	16011

+11,795 (1st intron)

SEQ ID NO:17

**The 5' (-5086 ) and 3' limits of the Human SM-MHC Promoter-Enhancer LacZ Transgene Tested in Transgenic Mice**

- 5 The number in the left margin refers to the position within an undefined BAC sequence contained in the public database (Accession # U91323 in GenBank). The start site (i.e. +1 position) of the SM-MHC gene corresponds to the BAC position 143,590.

10	-5086	TTTAAA ATTATTAAT CTTCTTTTT TTTTTTTGAA
	138541	GATGGAGCCT CTCTCTCTAG CCTAGGCTAG AGTGAATGG TGTGATCTTG GCTCACTGCA
	138601	ACCTCCACCT CCCAGGTTCA AGGGATTCTC CTGTCTCAGC CTCCAAGTA GCTGGGACTA
	138661	CAGGCCTGCA CAACCACACC CGACTAGTT TTGTATTTT AGTAGTGATA GGGTTTTACC
	138721	ATGTTGCCA GGCTGGTCTC GAACCTCTGA CCTCAAGTGT TCCCTCCACC TTGGCCTCCC
	138781	AAAGTGTGG GATTACAGGT GTGAGCCACT GTGCCCGGCC AAAAAATATT AAATCTTGAG
15	138841	GCACATGCAG GAGTAAGCCA TGCTCAGACC CAATCTTCGA TGTTACTAAA AATTGGAGGG
	138901	GATCACACTT CATGGTTTTG TTTTGTGTTG TTTTTTGAG ACAGGGCTT GCTCTGTTGC
	138961	CCAGGCTGGA GTGCACTGGT ACGATCACAG TTCACTGCA CCTCAAACTC TGGGGCTCAA
	139021	ACAATCCTCC TACTTCACTC TCTAGTTGGG ACTACAGGCA CACACTGCTG TGCTGACTA
	139081	ATTATTATTA TTATTATTAT TATTATTATT ATTATTATTA TTATTATTTT GTAGAGACAG
20	139141	GGATCTTGCT ATGTTACCTA GGCTGTTCTT GAACTCCTGG GCTCAAGCGA TCCTTCCGCT
	139201	GCAGCCTCTC AAAGTGCTAG GATTACAGGC ATGCCAGGCC ACTTTGGGGC TTTTTTAAGC
	139261	CAACAGCAAA AAAAGACTAT AAGAGAGAAA TTCCCCCTTG GCTGTTGTG TTCATGGATT
	139321	CGTGGAAACT CCCATTAAAC AGCCGGTCAC AGAAAAAGAT ATGCCAAGGA AAATTACTTG
	139381	ACAGCACTCA GTCAAAGTGA CATTAAAGA AGAGACTATT GCCTCCTCCA TCTTAAAAGA
25	139441	ACTGACCTT TGAGCCATGA GAAATGAAAC AGAGGCATCT GATCGAATGA TAACAATGCA
	139501	CTTCTGAAGA TTCAAACATC GGAACATTCA GCATTGGACA CATATCTATT GAATGACTCT
	139561	TAAGTGAACA TACTGTCCCT GCCTGCTTCC AGAGGGTACT AGAGAGGTCG GAGATGGTC
	139621	ATAAAGGCCA TCACATGTGC TGTCATATTT ACAATCAGA AAGGTACTTG AGGCAAAGAA
	139681	TCTGATCATC TTTGTTTTTC CTTGAGAAAA TGCGCTCAGA GAGGTTACT GACAATCCA
30	139741	AAGGTGCTTG GTGGTGTGTT AAGAGATCTG GGTTAAAC CTCAGACTGC TGTCTACTAT
	139801	GGCCTGTGTC AGAAAGACTG GGGTTGGAAT TCCTGTTCCA CCACTGCTGT GTTATTTAAC
	139861	CCCTCCAAAC CTAGATTCTC AACAAATAAA TGGGGTAGG GAGGAATTA AAGTATGTAC
	139921	CTTATTTTT AGAGACAACA TCTTGCTCTG TCGCCCAGGC TAGAGTGCAG TGGTGCAATC
	139981	ATAGTTCACT GTAGTCTCAA CCTTCCAAGC TCAAGAGATC CTCCTACCTC AGCCTCCCTA
35	140041	GTAGCTGGAA CTTCAGGCTA CACTACGCC AGCTGCTATT TATTATTTAT TTATTTATTG
	140101	AGATTGCATC TCACCATGTT GCCCAGGCTG GCTACTTAAA AAAAATTTT TTTTCAAGA
	140161	CAGGGTCTCA CTCTGCCACC CAGGCTGGAG TACAGTGACA GAGTCTCAGC TCACTGCAAC
	140221	CTCTGCCCTC CAGGCTCAAG TGATCTCCC ACCTCAGCCT CCCAAGGAGC TGGGATTACA
	140281	GGTACCCACC ACCACACATG GCTAACCTTT TATTTTTGT AGAGACAGGG TCTTGCTATG
40	140341	TTGCCCAAGG TGTTCTAAA CTCCTGAGCT CAAGCAATCC TCCTGCTTTG GCCTCCAAA
	140401	GTGCTAGGAT TACAGTTGTG AGCCACCATG CCTGGCTTG GCCACTTTAG TTTGCTTT
	140461	TTTTTTTTT TTTGAGTTGG AGTCTGCTC TGTCTCCAG GCTCCCAGGC TGGAGTGCAG
	140521	TGACACAATC TCAGCTCACT GCAACCTCTG CCTCCTGGGT TCAAGCAATT ATCCTGCTC
	140581	AGCCTCCCAA GTAGCTGGG CCACAGGTGT GCACCCACAT GCCCAGCTAA TTTTTATATT
45	140641	TTTAGTAGAA ATGGGGGTTT CACCATGTTG GCTAGGCTGG TCTTGAACCTT CTGACTTCAA
	140701	GTGATCCGCC TACCTTGCCC TCCCAAAGTG CTGGGATTAC AGGCAAGAGC CACCGTGCC
	140761	GGCTGCCATC TTTAATTTT AATAAAGGGT TGTTATATAA GGGTAGGTG AGAGAATGAA
	140821	GTAAAATGTA GTGTTACAGT CTCCAGTTGT TAATCACATT ATAATTATTC TCTTTAAAAA
	140881	GTTACCAACA AGTTATTTAA AGAATCGAAT GGAACCCCTT GGAAATACAG TGTTCATGCC
50	140941	TCTAGTATTA ATGCCAGTTT TTACTTCGAG GCCAGCAAGC TAGATTCGA TGGCCTTCCC
	141001	TTTCCAGGAT GGGAAAGCGGA TGATTGACTT CAATTTTCCC CCTCCCGTTA CTTCTGCT
	141061	CCACATCATT TCTGTGCTGA TGCAGGGACG ATTTCCACTC CTTTACAGC GTAGATGTTA
	141121	AAAGCCTGTG CGGAGCAGCT CATTCACTCAT TTCCGCAGA GCTTACCCC TCACTTCCCC
	141181	AGCCAGCTAA ATGCAGGCTG TTCTGACTC TCTGATCTAG GCCCATTGCA GGGTGAGGGC
55	141241	CAGGCTCAGG AGTTTCCAGG GTGAAAACCA GGTAAGCTTG ATGTTGGAAG GATGAAGAAG
	141301	GACCCAAAAG GGTCTGAGAT GCAGAGCTCT CCAGATGGGC CTGGGAGCCT GCAGGGAAAG
	141361	AGGCCTCTCT TTATATCCCG GAGGCCTGGT GCAACTCTAG TTGGTTTCAT GTTTGTTGCG
	141421	AGTAACAGCA GTCACATGA AGCGGTGCAC CATGTTCATT TTACATGGAT TCATCTCAAG

	141481	GAUTGCTTAC	AAAAAGGCCA	GGAAGTAGCT	GATGTTCTTC	CCATCTTACA	GGTAGGGAAA
	141541	TTGAGGCATG	GAGAGGCAAA	GTTACTTGCC	CATGGTCATA	TAGGTAGAAA	GCAGCACTGG
5	141601	CAGATTCAAA	GCCAGACATC	TACTCTCAGA	TACACGCCCT	GGGCCTCAAG	GCCAGTTGC
	141661	CTGGGCATTT	CCCTTTAATG	TCTCCTCTCT	GGAAAGTGAAT	GGTGTCACTCA	GAAAGGTTC
	141721	AGTGCCAGCA	CCAATCAATG	ACTGTCCCAG	TGAGAGCTTG	GTCAAATCCC	TTTACCCCTG
	141781	CAGGGACTCA	ATTTTCTCAC	CTGCAAAATG	GGGGTATTAA	TAAAGCCACC	CCCCGCACCC
	141841	CCGGCCCCCA	GCCCCCTCCAC	CTGGTTGCAA	GAGGAGTGGT	TGTAGACTAA	GGGCCTGCGT
	141901	CAAGTACAGA	ACCCAGGAGG	GGTCTGCCCA	ACTTTAACCC	TCTCTCCAAA	TCCTCTAGCC
10	141961	TGAAGCAGCA	AAAACCCACG	TGGGACTGGG	GGCTGCCCTT	TTCCGGGCCT	TCCCCAAGCA
	142021	GAGGGGTCCC	CATCTAGCCC	CGCGGGCAA	CGCGGGCCGG	TGGCTGCGTG	AAGGGCCCCC
	142081	TCCCCCGACG	CCGGGGAGCA	GGAAGGCCAC	TCGGCACCAT	ATTTAGTCAG	GGGGAGCCGG
	142141	CAGCCCAGAG	CTGGTATGCG	GCGCTGGGAA	TTCCTGCAGG	AAGGAGTCCG	CGCCTGCCCT
	142201	TTTTGGGTTG	TCTCCCGCCC	GCCGCTCCCG	CCGCTCCCGG	GGAGGGGGAC	CGGCCCCGCC
15	142261	CGGCCCCGCC	CGGGAACCTC	GGAGGAGCTG	GTGCGCGCG	GGGAGCGGAG	CGCCCGGGCT
	142321	GCCCCGGGGT	CCCCGGCCTG	GCGCGGGGCC	AGCCCACCCG	CTCGACTTCC	TTTTATGCC
	142381	TGTGTGTGCG	TGCGTGGACA	GGAGCGGGGA	GGGAGGGACG	GGGAGAAGAC	GGAGAGCTG
	142441	GGGAAGAGAG	AGAGAGAAAG	CGCAGAGATA	GGAGTGGAGAC	ACGCGGGAGA	GATGGAGAGC
	142501	AAGAGACACA	GAGACCAGAG	ACAAAGTGG	ACAGGAGGGA	GAGACAGATA	CATCGACAGA
	142561	TCTAGAGAAG	CGAGAGGGAC	AGAGACAAAA	GATAGAGCGA	GAGACAGCAA	TGATCAGAGT
20	142621	GACAGACATG	CAGAGACAGT	GGCAGAGACA	GAGCGAGAGA	GCCTGTGATG	GAGAGAGACA
	142681	GGGAATGCAA	TTTTAGGCGA	GGAATCCTTG	GGGAAGGGAA	TTTGTGAAAG	GGAACCTCGA
	142741	GAUTCTGGGG	GCACACCCAC	TTTCTCCTTG	GATCTGACA	CTTGATCTT	GTAAATAACG
	142801	TAATTATCAC	CGCCACCGCC	TTCCCCCATT	TTGTAGCTAT	GGACACCAAAG	TCTCAGAGAA
	142861	GTGAAGTGC	TTGCCCAAGG	TCACGCAGCT	GGCGAGTGGC	GCACAGGGGA	GGGGGACAGC
25	142921	TGAAATAATC	ACAGTGGGCT	TATTTTTAAT	TTTTATTTGT	ATTTTGGTCG	TGGTGTATGT
	142981	GGTGGAGGTG	GAGATGGCAA	GTGGGGAAAA	GTAAAAAACTT	CCCCTTCCTG	CACGGTTCCC
	143041	AGCAAGGGTG	GGGGCCTCCT	GTCTTGCACT	TTGCAAAGTT	CAAGAAATCC	CCTTCCCTA
	143101	CCCTTCACGC	TGACACAGCCG	GCCCTCTTTC	CAGACAGTGC	GATGCCAATA	AAATGGGAAG
	143161	TGGGGTGGGA	GATGTCAAGT	CAGATCCACC	ACAGCCCCGA	CACGGGGAGG	AAGAGGTTAA
30	143221	AGCCTTGC	GCCGGAACCG	ACTCAGGGAA	GACGTTCTCA	AGCATCCCGC	ACAGACACTG
	143281	CCTGCTCGAC	CCCCTTTCTC	TAGGGATCCG	GAGCGTCTGC	GACCGCCTGG	GGCGGGGGCT
	143341	GAGACTCCCG	TCCCTGTGCG	CACCTGTTCC	GTGCGCCCTT	GTGCGGTGCG	CACCTGTTCC
	143401	GTGCACCCCT	GTCCCAGCG	CCCCAGCTCC	TTGCGCTCCC	GCCGGGGGTG	CGCCCTGAG
	143461	GGGGCGCGGC	GAGGGGGCCG	CGAGGGACCC	TCCCCAACTC	CACCCCTTCG	GCCTCTCCCC
35	143521	CTTCCCAGC	CGCGGGCAGC	TCCGGGTCTA	TAAGAGAGGG	CGTCCGAGGA	CGCGCAGGGA
	<b>TRANSCRIPTION START SITE +1</b>						
	143581	GATTTGGACG	<u>CTCCGGCCTG</u>	GGAGGTGCGT	CAGATCCGAG	CTCGCCATCC	AGTTTCCCTCT
	143641	CCACTAGTCC	CCCCAGTTGG	AGATCTGTAA	GTAGTAGTTG	TCATTCTGGG	GGCAGATTGC
40	143701	AGGGCAGGGG	GGTGTAAAAA	GTCCTATAGG	GTATTCTATA	GGGGCTGGGG	TGCACTTACG
	143761	GGTCCCTGTT	GTCAACCTCG	TAAGGGCCAT	GGTGGGGGCA	GAGTTGTGAT	TTGGATCTCT
	143821	CTCTGCCTTA	TCGTCTTAGA	TTATCCTAGA	CTTTCCTCAA	ACAGCATTTC	TTAAGATTGC
	143881	CAGTGGAGAG	TACCATTTG	GGGGTGCTTA	TTAACGATAT	CAATGCCTGG	ACCCAACTCC
	143941	ATTTCCCAAC	TCTAGAATCC	CCAGAAAAAC	TGCCTTAAAAA	AAAAAAAAT	TAGTCCCGAG
	144001	TGATTCTGT	TAAGAGGCTA	ATCCAGGAGA	TATGCTCCCT	TGGAAATCTC	AGAGGTCCGG
45	144061	TGCAGACAAT	CAAGGCATCT	CACTTTTATT	CTAGGCACCA	AAAAAATTAC	AGCTGAACCT
	144121	CACTGAAAAG	TCACTTGCTA	TCACACAGAA	GGGCAAAGTG	AGGCTCCTTG	TGGATTGAC
	144181	CGTATTGCA	AGTGTGTTG	ATAATGCATT	AAATCAGTTA	AAAACACATG	GGCATAGGCT
	144241	TAGCAGAAAAG	GAGTGTGTT	GTTTTTTTT	TTAACATCAGT	TTAGGGGAGG	TTCTTCTATG
	144301	TTGAGAACCC	CTGGGAGATA	AGGCTGGTTG	TGATCTAGTT	TGTTACAGCC	CACTTTTCC
50	144361	TCTTCTCCAA	ATTAaaaaaaa	AAAAAAACAA	CTCACCCAGG	TTGACCCCAA	AGGGCCCCAA
	144421	GATAACCCAGG	TGGGCTCCAA	AGTCTCATT	TGCTTCCACG	ATCTGCAGGT	GGGTTAGGTA
	144481	AGATTACACT	AGAATTCCC	GCAGAGCCAC	CTGTGTCAAT	GCCACTCTCG	TGCCCAACCA
	144541	AATGGGTAAA	ACGAGAGAAA	GTGTGGCTAC	TGCTGTGTTGT	AAGTTTCTT	CCAGCACAGG
	144601	GTCTGGTAGG	GATTTGCCA	CTTGAGAAAA	GGTACCATCC	AAAGCCATGC	TTGTCAAGAA
55	144661	GTAAAAGAAA	ATATTTAGAA	ACCCAAAGGTG	GGAGTGTTTA	GTTGCAGTAT	GAAGAACTGA
	144721	GAGATTAAAT	GGTGAACGTG	CCGTCCGGGG	TTTGGCAAAA	AGAATGCAGG	CTATTAATAA
	144781	ACTGCTTGC	ATAGTTTTT	GTTCCTTGA	TTTACTCAAC	GATACTATT	TAGAATTGTT
	144841	CAGAGACGGA	ACTGACGCT	GAACTGAAAG	TCATTAGGTG	GCAGGGTGTG	AAATAAGATA
	144901	GAGAATTTCG	TTTGAAGGAA	ATTGATGTTT	TCCCTTTGAG	ATAGCTACCG	TTGATGGAAC

144961	ACTTCAGTGC	CACATGCTGT	TGCAACATT	AACTTAATT	ATCTCATT	ATCTTGCAA
145021	CAACTTCATA	AGAAAGGCTT	TATGATGCCT	GTTTAGTATA	CAAGGCAGCT	GAGGCTCAGA
145081	GAGGTAAGT	GTCACACAGC	CAGCAAGTGG	TAGAACCCAT	TCCCCGGTCA	GTTTGAGTCC
145141	AAGTTCATAC	CCTTGACCCC	ACTATCTTC	TTCTTACCA	TGGACACAAA	CTTGTGAGGG
145201	TCAGGTTCT	GGTGGGACTA	AATGCTTCCA	ACAAAGTAAA	TGTTTATCAC	CGTGTCCCTT
145261	GAAGAAAACA	TAAGTGA	TTTGACAT	TTAAAATAAA	AGGCAGTGT	TGTCCCCTGA
145321	TTGAGGGG	GACCTAGCTG	AAACCAGTGA	CCCTAGGTGG	GCTGCCATGC	CGAGAGTCCA
145381	GAACGTGAAC	TAGCTGGTC	TTTCCGAGA	AGCCGCCAGG	CTTGCTTGT	AAACACCATG
145441	TTTTTTATT	ATCATGTCCG	AAATAGATGT	GTTATTCCGT	ACAAGGTATC	TGTTATGGAT
10	145501	TTGTTATCAT	TACTTTCCG	TGGGAGGGCA	GAGATTGAGG	CAAACATGCC
145561	AGCGTTTCC	ATGAGGCCAT	CCCCGGCCCC	CTCGTCAGTT	ACCCAGCCTT	GCACCGCAGC
145621	CCGGTTGGTC	CTGGCCCTGG	GGATTGTCT	ACCATGTCCC	TCACCCATTG	AAGAACTAGT
145681	GGAGAAACCC	TAAGGAGAAG	AGATTGGGA	GGAAAGTGGG	ATTCTTTTT	CCTACCCCCCT
145741	CTTATTCA	GGTTGATTT	TTTGGGTGG	GGGGTGGGAG	GGAAATTGTCT	CCTTCCACA
15	145801	GGTCTGAAT	CCAAACAGGT	GGGTCTTCA	CGTTAGGCAC	AAGCGTGTAA
145861	AGATATATAG	TAGATTTTC	TTGAAAACCA	AGTTCAATAT	TCAATCCAGT	AGAACATAG
145921	AAGGCCATAA	GCAAATTAA	AAATCATCTC	CCGCACCTCC	CCAAACCTCA	CTTTCTCATC
145981	CGGGAAATGG	GGCTAATGAG	AATAACTCAT	GTTTTTGAGG	CACTTTGCC	TGGCGAGATG
146041	CTAACGCTT	TGTGGACATT	ATCTTACGT	TTCATAACAA	CCCTTAGAG	TAGATACTGT
20	146101	TATTCTA	GGCTTTATT	TACACATATG	GAGTCTGAAT	AACTTGCTTA
146161	GCTAACCA	AGGAAAAGA	AGATTCTACA	AATCTAGGT	TTTCTAACTC	CAGAGTTCTA
146221	CAGATTACCC	TCATGGGAGG	ATTGATGAG	CTAATGTGT	TGAAGGGTTT	AGCACAGTGC
146281	CTGGCCCTG	GTAAGCTCA	GTGATGGTT	TTTATAGCAA	ACACAAACCAG	AGAGTTCAAG
146341	ATGTTGCTC	AGTATGGCAT	GGCTCATCTT	TGGCAGAAC	GGGAAGCTA	AACTATGTGG
25	146401	CCGTTAAAGG	AGAACCTTCT	CTTAATTTC	TTCCCTTGA	TCTCATAAAC
146461	TTGGGCTGA	AAGTGGTGAT	TAGAATCTT	AATATATTAA	GCTACCATT	CTTACCTGG
146521	TTGGGAATGT	TACAAATTCC	AATTACATT	GTTTAGGGTT	TTGTTTGT	GTTTTGAGA
146581	CAGAGTCTG	CTCTGTCGCC	CAGGCTGGAG	TGCAGTGGT	CGATCTGGC	TCACTGCAAC
146641	CTCCGCC	TAGGTTCA	CACTCTCCA	GCCTCAGCCT	CCTGAGTGA	GAGTAGCTGG
30	146701	GTTTATAGGC	GCCCCACCACC	ATGCCTGGCT	AATTTTTGT	ATTTTAGTA
146761	TTCACCATAT	TGGCCAGGCT	GGTCTCGAAC	CGCTGACCTC	AAGTATTG	CCTGGCTTGG
146821	TCTCCAAAG	TGCTCAGATT	ACAGGCGTA	GCCACCGCGC	CTGGCTTATT	TAGGGCTTGG
146881	ATGGCATACT	TTAAGGGATG	GCCTTTTG	TCTCTAGGT	TTCTCCTTCC	ACTCCTGACC
146941	TTCAACTT	TAACCCCTGGC	CACACATGG	AGGAAAGACT	GAATTAGAG	AAAGGCAGGC
35	147001	AAGAATTG	AAAGAACCTT	GTATGTGATC	CAAGGACAGA	GGAAAGAAGCT
147061	GCTGAAAGGG	GAGGTCGGAC	ATCTGTGACT	TGTATCAGGG	TTTCAGGGC	TAAGGAGGAA
147121	CAACCTCATC	AAAGTTGCTA	GGAAAGGGCC	ATAGAGGCCA	GGTATGGCAG	GTCATACCTG
147181	TAATCCCAGC	AATTGGGAG	GCTGAGGTGG	GGGGATGGCT	TGAAGTCAGG	AGTTGAGAC
147241	CAGAGTGGC	AAACATAGCGA	GGCACCATCT	CTACAAAAAA	ATTTTAAAAA	TGAGCTGGC
40	147301	ATGGTGGCAT	GCATCTGTAG	TCCTAGTT	TCAGGAGGT	GAGTGGAGCA
147361	TTGAGCC	GAGTCAGG	CTGCCGTGG	CCCTGATTG	ATCACTGTT	TCTAGCCTGG
147421	GCAACAGAGT	GAGACTCTGT	CTCAAAAAAA	AGGTGAGGGG	CATAGAACTT	TACTGTACCA
147481	GGCTGAAAAA	TACAAGGCC	AGAGAGGGCA	AGTGACTTGC	CTAGCATCAC	CCAGCGAGTT
45	147541	TTGGGAGAG	CTGAGACTTG	TAACTCGAAG	ACCTAAGGAT	CTTCCACAGG
147601	GCTTGT	GCTCAAGGG	TGAAGCAGT	AGTTGTTAGG	ACAGGACTGT	GAATAGGGCT
147661	GACATATTCA	GATGTGTAA	ACATCGCTAA	TGCCATCTCT	GAGTAAATT	GGCTCAAAC
147721	AGATCGGGAT	TCTAATCTG	GTTCCCCAAC	TTTGCAAGG	GAGGGCCTTG	CATTTACCTT
147781	TCAAGACCC	GATAGGCTT	GCAGGAAAAT	GGGAATAATA	GATAATGCCA	CTCTTCATC
50	147841	CTTGGACTT	TTGTCTAATT	ATATGAATT	ATCTGTAGGA	TAAATTCCC
147901	TGCTGAGTT	AAGGGCATGC	GTATCTAAA	TTAATAGATA	TTGCAAATGA	CTGGCTAAAG
147961	ACATTGCA	CCAGGTGCAG	TGGCTCACGC	CTGTAATCCC	AGCACTTTGG	GAGGCCGAG
148021	CAGGTGGTC	ACCTGAGGT	AGGAGTCAA	GACCAGCCTG	GCCAACATGG	TCTCTGCTAA
148081	ACCCTATCTC	TACTAAAT	ACAAAAATT	TCTGGCATG	GTCGTGGGCA	CCTGTAATCC
148141	CAGCTACTCG	GGAGGCTGAG	GCACGAGAAT	CGCTTGAGCC	TCAGAGGCAG	AGGTTGCATT
55	148201	GAGCCGAGAT	CACACCCTG	CACTCCAGCC	TGGGCAAAGA	GTGAGACTCG
148261	AAAAAAAAAA	AAGGCATTG	AAATTGCAAC	TTGTTGCAGT	CACATATGAC	AGCAGTCCCC
148321	ATCCTCTGG	CACCAAGAGAC	TGGTTTGT	GAAGACAATA	TTTCCAGGG	TGGAGTGGGG
148381	AGGATGGTT	TGGGATGAAA	CTGTCCCACC	TCATCATCAG	GCATTGGTTA	GATTCTCATC
148441	AGGAACGTAC	AACCTAGATC	CCTTGCAGGT	GGAGTTGGCA	ATAGGGTTTG	TGCTTCTGTG

	148501	AAAATCTAAT	GCTGCTTATC	TGACAGGAGG	CGGAGCTTAG	GCAGTGATGG	TCACTCACCC
5	148561	ACCGTCCCCT	CCTGCTATGT	GGCCTGGTC	CTAACAGGCC	ATTGACTGAT	ACTGCAGCAC
	148621	AAGGGTGGG	GACCCCTGAC	ATAGGAGACT	ATACATTAT	TTAAAGCTGT	GGTATGCCAG
	148681	AATTGTAAAA	TATAAAACAC	AGTGGGGCTT	TTAGGGCCAG	AAATAATCAG	TTCTTGCTG
	148741	CTTCCAGAAG	CATCCTTCAC	AGGGGCTACC	GTAACTCTT	CCAACCAAGT	TCTCTTGTT
10	148801	GGGAGGAAAA	AATAGTGT	TGCATTAAGA	GAACCTCTT	CTGGAGTTAC	TTGAAACCAT
	148861	TGGTATTTCAG	ATGATTAGGC	AGATGTCACA	AGGCAATAAG	AATGTGACAG	GTTCACCAT
	148921	CACTTTTTT	CCTGAAAAG	TGAAGTAGGG	CTTCTTGGG	AAACAAGCC	TGGGAGGTGG
	148981	GGGGATGTGA	ATGGTGAGGG	GAGGGTAGAA	ATGGTGGAGT	AGGGTCAGGG	GCAAGAAAGG
15	149041	GACTTTCTGC	TAAGAATTAA	TCGGGTGTCC	ATTACTCTT	AGCAGAAAAC	TAGGATTAGA
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	149161	GACCTCGGTC	ATCTTAGCAG	GAAAATAGCA	ATAGCAGGTG	ATGCCACCTT	ACAGAGCGCT
	149221	TAGGAGACAG	TGAGATGGTC	TATATAGGAA	GCTGTCTGGC	CTGATACCTG	ATGAATACAA
	149281	GGGGCCCAAT	AAATACAGTG	GCTGTTATGA	ATAATAGATC	AAAATGCCT	TTTTGGTACT
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	149461	CAGCTGGCGA	ATTGTGTTGC	AGATCAGGTT	CAGAGAACTT	CTGTTTGCC	TGTGTGGCAT
	149521	TCATTCAATC	GT	TTTATTGAA	AATAGAGATG	GGATCTCACT	GTGCTGCCA
	149581	GAGCTCCTAA	TTCAAGCAAT	CCTCTGGCT	TGGCCTCCCC	TAGTTCTTGG	ATTACAGGTG
25	149641	TGAACCAC	TATCCAGCCC	TTTATGACAT	TTAGAATATG	AGCAATTTT	CTTTTTCTT
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	149761	CTTGGCTCAC	TGCAACCTCT	ACCTCCCAGG	CTCAAGCGAT	CTTCCCACCT	CAGCCTCCG
	149821	AGTAGCTGGG	ACTACCGGCA	TGTGCTGCCA	TGCCTGGCTA	ATTTTTGTAT	TTTCTGTAGA
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	150061	ATCTCCAAA	ACATTTGGGT	TTTTGTCTCT	GGTCCAAAAT	CTTTAGCAA	TGGCTTGCA
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	150481	AGGCATCAGC	CACCATGCC	GGCCTAATT	ACTTTTATT	AATGCTGAAG	CAGAGAGGGC
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	150661	CCCATAAAAT	GGTAGAGCTG	GGCCAGCCTA	CCATTGATT	ATTTCC	AATGAAAAT
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45	150841	TCAAAAATG	AAGCAGCCTG	GGCGCGGTG	CTCATGCTT	TAATCCTAGC	ACTTTGGGAG
	150901	GCTGAGGCAG	GCAGATCGCT	TGAGCTCAGG	AGTTTGAGAC	CAGCCTGGG	CACAAAGTGA
	150961	GACCCCTGTC	TCTACAAA	AATGCAAGAA	TTAAAAAATT	AGCTGGGTGT	TCTGGTGC
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	151381	GACCCCATC	TCTACAAA	ATAAGCCGGG	CATAGTGGCC	CACACCTGAG	GTGGGAGGAT
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	151501	TGGGCA	TATA	GTGAGACCC	TCCCCCAACC	AAAACATTG	AGAGCAGCTC
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	151621	GGAGCTGATG	CAGGACCTTA	AACATGAGCG	ATGGTGGAGG	AGGGAGGGTT	GGGAAGGTC
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55	151741	GTGTGTTCA	CTTGGCCCCG	TCTGATCTTC	TGCAGTTGGT	ATTCCGAGTT	GAGTTTGACT
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	151861	AAGCTGTTG	TCTCAACTGG	AAGCAGTTT	GCCACCC	GGGACATCTA	GCAGTGTG
	151921	GAGACATTT	TGATTGTCA	GAGTGGAGGA	AGGGTGTCA	CTGGCATCAG	GTGGGCAAGAG
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